```
logo
NTDB id 1127 NGFG RS10560 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMF....VLFC....MLFSCAFSLAAVNINAASQQEL.EALPGIGP   71
NTDB id 1126 NGFG RS09375 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMF....VLFC....MLFSCAFSLAAVNINAASQQEL.EALPGIGP   71
NTDB id 1125 NGFG RS07215 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMF....VLFC....MLFSCAFSLAAVNINAASQQEL.EALPGIGP   71
NTDB id 1108 NGFG RS06525 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMF....VLFC....MLFSCAFSLAAVNINAASQQEL.EALPGIGP   71
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    ......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAAQ.............VEEGIVITVNINTASAEELATLLKGIGL   61
NTDB id 1150 A1552VC RS08580 WP 001166105.1   ......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAAQ.............VEEGIVITVNINTASAEELATLLKGIGL   61
NTDB id 1395 DSB67 RS04445 WP 010649155.1     .............MKWMLTLCLLILAPMSWAE.TKTKAD.............KYEGIEITVNVNSATAQEIATLLIGIGE   53
NTDB id 973135 AABD50 RS04485 WP 023624118.1  .............MKWILTLCLVVFAPLSLAA..DTKAD.............KYEGIEITVNINTASAEEIATMLKGIGE   52
NTDB id 1294 VP RS04495 WP 021449594.1        .............MKWILTLCLVVFAPLSLAA..DTKAD.............KYEGIEITVNINTASAEEIATMLKGIGE   52
NTDB id 1362 HI 1008 AAC22669.1               ........MKTL..FTSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEIQKSLTGIGA   70
NTDB id 1382 A4U84 RS07375 WP 062924234.1     ........MKKLSRYTFSLLFG.LFSATTFAEEKVTTSTEHTPVVMQQ.VAATQKVNANVVNINTATASEIQDKLVGIGA   70
consensus                                              * **** ** ******* *  *  *                     **  *!*!*!** !*   ! !!!

logo
NTDB id 1127 NGFG RS10560 WP 003703428.1      AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1      AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1      AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1      AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    KKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   KKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1395 DSB67 RS04445 WP 010649155.1     KKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL...................    95
NTDB id 973135 AABD50 RS04485 WP 023624118.1  KKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEARILL...................    94
NTDB id 1294 VP RS04495 WP 021449594.1        KKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL...................    94
NTDB id 1362 HI 1008 AAC22669.1               KKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF...................   112
NTDB id 1382 A4U84 RS07375 WP 062924234.1     KKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE..................   113
consensus                                     *!!**!**!!  *! !*****!* !*!!! !** ********
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X non conserved

X similar

X ≥ 50% conserved